1600

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/500,746

DATE: 10/18/2001
TIME: 09:45:14

Input Set : **A:\ES.txt**
Output Set: **N:\CRF3\10182001\I500746.raw**

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  3 <110> APPLICANT: Winchester , Robert J.
  4         Gulko, Percio
  5         Seki, Tetsunori
  7 <120> TITLE OF INVENTION: USES OF INHIBITORS FOR THE ACTIVATION OF CXCR4 RECEPTOR BY
SDF-1 IN
  8         TREATING RHEVMATOID ARTHRITIS
 10 <130> FILE REFERENCE: 0575/57005-B
 12 <140> CURRENT APPLICATION NUMBER: 09/500,746
 13 <141> CURRENT FILING DATE: 2000-02-09
 15 <160> NUMBER OF SEQ ID NOS: 23
 17 <170> SOFTWARE: PatentIn version 3.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 13
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Artificial Sequence
 24 <220> FEATURE:
 25 <223> OTHER INFORMATION: Primer
 27 <220> FEATURE:
 28 <221> NAME/KEY: misc_feature
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 39 <213> ORGANISM: Artificial Sequence
 41 <220> FEATURE:
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 54 <211> LENGTH: 24
 55 <212> TYPE: DNA
 56 <213> ORGANISM: Artificial Sequence
 58 <220> FEATURE:
 59 <223> OTHER INFORMATION: Primer
 61 <220> FEATURE:
 62 <221> NAME/KEY: misc_feature
 63 <223> OTHER INFORMATION: Primer
 66 <400> SEQUENCE: 3
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 70 <210> SEQ ID NO: 4
 71 <211> LENGTH: 12
 72 <212> TYPE: DNA
 73 <213> ORGANISM: Artificial Sequence
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75 <220> FEATURE:
76 <223> OTHER INFORMATION: Primer
78 <220> FEATURE:
79 <221> NAME/KEY: misc_feature
80 <223> OTHER INFORMATION: Primer
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88 <211> LENGTH: 24
89 <212> TYPE: DNA
90 <213> ORGANISM: Artificial Sequence
92 <220> FEATURE:
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95 <220> FEATURE:
96 <221> NAME/KEY: misc_feature
97 <223> OTHER INFORMATION: Primer
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106 <212> TYPE: DNA
107 <213> ORGANISM: Artificial Sequence
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112 <220> FEATURE:
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114 <223> OTHER INFORMATION: Primer
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121 <210> SEQ ID NO: 7
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123 <212> TYPE: DNA
124 <213> ORGANISM: Artificial Sequence
126 <220> FEATURE:
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129 <220> FEATURE:
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131 <223> OTHER INFORMATION: Primer
134 <400> SEQUENCE: 7
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139 <211> LENGTH: 12
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141 <213> ORGANISM: Artificial Sequence
143 <220> FEATURE:
144 <223> OTHER INFORMATION: Primer
146 <220> FEATURE:
147 <221> NAME/KEY: misc_feature
148 <223> OTHER INFORMATION: Primer

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158 <213> ORGANISM: mouse
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162 <222> LOCATION: (337)..(337)
163 <223> OTHER INFORMATION: x= to any amino acid
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168 <222> LOCATION: (376)..(376)
169 <223> OTHER INFORMATION: x= to any amino acid
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179             20                  25                  30
182 Gln Gly Arg Ile Pro Tyr Pro Arg Ser Ala Val Cys Val Tyr His Leu
183         35                  40                  45
186 Ser Asp Ile Gln Thr Val Phe Asn Gly Pro Phe Ala His Lys Glu Gly
187     50                  55                  60
190 Pro Asn His Gln Leu Ile Ser Tyr Gln Gly Arg Ile Pro Tyr Pro Arg
191 65                  70                  75                  80
194 Ser Ala Val Cys Val Tyr Ser Met Ala Asp Ile Arg Met Val Phe Asn
195                 85                  90                  95
198 Gly Pro Phe Ala His Lys Glu Gly Pro Asn Tyr Gln Trp Met Pro Phe
199             100                 105                 110
202 Ser Gly Lys Met Pro Tyr Pro Arg Ser Ala Val Cys Val Tyr Ser Met
203         115                 120                 125
206 Asn Asp Val Arg Arg Ala Phe Leu Gly Pro Phe Ala His Lys Glu Gly
207     130                 135                 140
210 Pro Met His Gln Trp Val Ser Tyr Gln Gly Arg Val Pro Tyr Pro Arg
211 145                 150                 155                 160
214 Ser Ala Val Cys Met Tyr Ser Met Ser Asp Val Arg Arg Val Arg Arg
215                 165                 170                 175
218 Val Phe Leu Gly Pro Tyr Ala His Arg Asp Gly Pro Asn Tyr Gln Trp
219             180                 185                 190
222 Val Pro Tyr Gln Gly Arg Val Pro Tyr Pro Arg Pro Gly Thr Cys Pro
223         195                 200                 205
226 Gly Gly Ala Phe Thr Pro Asn Met Arg Thr Thr Lys Asp Phe Pro Asp
227     210                 215                 220
230 Asp Val Val Thr Phe Ile Arg Asn His Pro Leu Met Tyr Asn Ser Ile
231 225                 230                 235                 240
234 Ser Pro Ile Pro Gly Thr Cys Pro Gly Gly Ala Leu Thr Pro Asn Met
235             245                 250                 255
238 Arg Thr Thr Lys Glu Phe Pro Asp Asp Val Val Thr Phe Ile Arg Asn
239             260                 265                 270
```

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242 His Pro Leu Met Tyr Asn Ser Ile Tyr Pro Ile Pro Gly Thr Cys Pro
243         275                 280                 285
246 Gly Gly Thr Phe Thr Pro Ser Met Lys Ser Thr Lys Asp Tyr Pro Asp
247     290                 295                 300
250 Glu Val Ile Asn Phe Met Arg Ser His Pro Leu Met Tyr Gln Ala Val
251 305                 310                 315                 320
254 Tyr Pro Leu Pro Gly Met Cys Pro Ser Lys Thr Phe Gly Thr Phe Ser
255             325                 330                 335
W--> 258 Xaa Ser Thr Lys Asp Phe Pro Asp Asp Val Ile Phe Ala Arg Asn His
259         340                 345                 350
262 Pro Leu Met Tyr Asn Ser Val Leu Pro Thr Pro Gly Thr Cys Pro Ser
263         355                 360                 365
W--> 266 Lys Thr Phe Gly Gly Phe Asp Xaa Ser Thr Lys Asp Leu Pro Asp Asp
267     370                 375                 380
270 Val Ile Thr Phe Ala Arg Ser His Pro Ala Met Tyr Asn Pro Val Phe
271 385                 390                 395                 400
274 Pro Met His Arg Arg Pro Leu Ile Val Arg Ile Gly Thr Asp Tyr Lys
275             405                 410                 415
278 Tyr Thr Lys Ile Ala Val Asp His Lys Arg Pro Leu Ile Val Arg Ile
279             420                 425                 430
282 Gly Thr Asp Tyr Lys Tyr Thr Lys Ile Ala Val Asp Gln Arg Arg Pro
283         435                 440                 445
286 Leu Val Val Arg Thr Gly Ala Pro Tyr Arg Leu Thr Thr Ile Ala Val
287     450                 455                 460
290 Asp Gly Gly Arg Pro Leu Phe Leu Gln Val Gly Ala Asn Tyr Thr Phe
291 465                 470                 475                 480
294 Thr Gln Ile Ala Ala Asp Asn Asn Arg Pro Ile Val Ile Lys Thr Asp
295             485                 490                 495
298 Val Asn Tyr Gln Phe Thr Gln Ile Val Val Asp
299         500                 505
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303 <211> LENGTH: 396
304 <212> TYPE: PRT
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307 <400> SEQUENCE: 10
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313 His Met Phe Glu Asn Glu Ile Ser His Arg Thr Gly Ser Trp Asn Phe
314             20                 25                 30
317 Ala Pro Asn Pro Asp Lys Gln Trp Leu Leu Gln Arg Thr Ser His Ala
318         35                 40                 45
321 Ala Pro His Gly Pro Glu Asp Ser Ala Pro Gln Phe Ser Glu Leu Tyr
322     50                 55                 60
325 Pro Asn Ala Ser Gln His Ile Thr Pro Ser Tyr Asn Tyr Ala Pro Asn
326 65                 70                 75                 80
329 Met Asp Lys His Trp Ile Met Gln Tyr Thr Ala Thr Pro Ala Pro His
330             85                 90                 95
333 Ser Pro Trp Thr Ala Ala Pro Gln Tyr Gln Lys Ala Phe Gln Asn Val
334             100                 105                 110
```

```
337 Phe Ala Pro Arg Asn Lys Asn Phe Asn Ile His Gly Thr Asn Lys His
338         115             120             125
341 Trp Leu Ile Arg Gln Ala Lys Gly Lys Met Asn Asp Val His Ile Ser
342     130             135             140
345 Phe Thr Asp Leu Leu His Arg Arg Arg Leu Gln Thr Leu Gln Ser Val
346 145             150             155             160
349 Asp Glu Gly Ile Glu Arg Leu Phe Asn Leu Leu Arg Glu Leu Asn Gln
350             165             170             175
353 Leu Trp Asn Thr Gly Pro Met Leu Pro Ile His Met Glu Phe Thr Asn
354             180             185             190
357 Ile Leu Gln Arg Lys Arg Leu Gln Thr Leu Met Ser Val Asp Asp Ser
358             195             200             205
361 Val Glu Arg Leu Tyr Asn Met Leu Val Glu Thr Gly Glu Leu Glu Asn
362     210             215             220
365 Thr Thr Pro Met Thr Asn Ser Ser Ile Gln Phe Leu Asp Asn Ala Phe
366 225             230             235             240
369 Arg Lys Arg Trp Gln Thr Leu Leu Ser Val Asp Asp Leu Val Glu Lys
370             245             250             255
373 Leu Val Lys Arg Leu Glu Phe Thr Gly Glu Leu Asn Asn Thr Tyr Ala
374             260             265             270
377 Ile Tyr Thr Ser Asp His Gly Tyr His Leu Gly Gln Phe Gly Leu Leu
378             275             280             285
381 Lys Gly Lys Asn Met Pro Tyr Glu Phe Asp Ile Arg Val Pro Phe Phe
382     290             295             300
385 Met Arg Gly Pro Gly Ile Pro Arg Tyr Ile Ile Tyr Thr Ala Asp His
386 305             310             315             320
389 Gly Tyr His Ile Gly Gln Phe Gly Leu Val Lys Gly Lys Ser Met Pro
390             325             330             335
393 Tyr Asp Phe Asp Ile Arg Val Pro Phe Phe Ile Arg Gly Pro Ser Val
394             340             345             350
397 Glu Pro Tyr Ile Phe Tyr Thr Ser Asp Asn Gly Tyr His Thr Gly Gln
398             355             360             365
401 Phe Ser Leu Pro Ile Asp Lys Arg Gln Leu Tyr Glu Phe Asp Ile Lys
402     370             375             380
405 Val Pro Leu Leu Val Arg Gly Pro Gly Ile Lys Pro
406 385             390             395
409 <210> SEQ ID NO: 11
410 <211> LENGTH: 102
411 <212> TYPE: PRT
412 <213> ORGANISM: Human
414 <400> SEQUENCE: 11
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417 1           5               10              15
420 Asn Gly Pro Phe Ala His Lys Glu Gly Pro Asn Tyr Gln Trp Met Pro
421             20              25              30
424 Phe Ser Gly Lys Met Pro Tyr Pro Arg Pro Gly Thr Cys Pro Gly Gly
425         35              40              45
428 Thr Phe Thr Pro Ser Met Lys Ser Thr Lys Asx Tyr Pro Asp Glu Val
429     50              55              60
```

VERIFICATION SUMMARY
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L:258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:469 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:510 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:1240 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:1244 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:1256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:1340 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
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